```
(FILE 'HOME' ENTERED AT 15:04:47 ON 11 JUL 2003)

      FILE 'BIOSIS, SCISEARCH, MEDLINE' ENTERED AT 15:04:54 ON 11 JUL 2003
               ACTIVATE DONGMEI/L
               ---------
L1               STR
L2    (          1) SEA FILE=REGISTRY SSS SAM L1
L3    (         38) SEA FILE=REGISTRY SSS FUL L1
L4               STR
L5    (         29) SEA FILE=REGISTRY SSS SAM L4
L6               STR
L7    (         29) SEA FILE=REGISTRY SSS SAM L6
L8    (        473) SEA FILE=REGISTRY SSS FUL L6
L9    (        705) SEA FILE=CAPLUS ABB=ON  PLU=ON  L8/BIOL OR PROC/PROC
               ---------
L10        1461 S ((FENG, D?)OR(FENG D?))/AU
L11          20 S L10 AND PROSTATE CANCER
L12          20 S L11 AND CONJUG?
L13          10 DUP REMO L12 (10 DUPLICATES REMOVED)
L14           2 S L13 AND VINBLAST?
L15       73223 S PROSTATE CANCER
L16         319 S L15 AND VINBLAS?
L17          70 S L16 AND CYTOTOXIC
L18          38 DUP REMO L17 (32 DUPLICATES REMOVED)
L19           3 S L18 AND CONJUGAT?
```
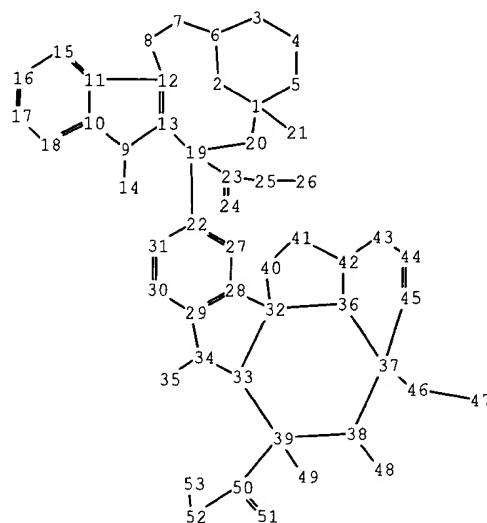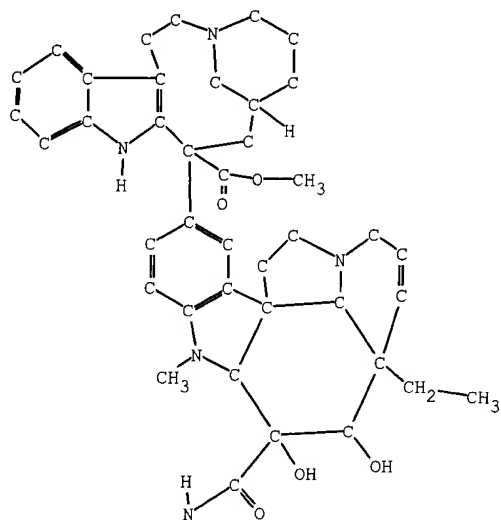
```
L1                STR
L2   (          1)SEA FILE=REGISTRY SSS SAM L1
L3   (         38)SEA FILE=REGISTRY SSS FUL L1
L4                STR
L5   (         29)SEA FILE=REGISTRY SSS SAM L4
L6                STR
L7   (         29)SEA FILE=REGISTRY SSS SAM L6
L8  (        473)SEA FILE=REGISTRY SSS FUL L6
L9  (        705)SEA FILE=CAPLUS ABB=ON  PLU=ON  L8/BIOL OR PROC/PRO
```

chain nodes :
    14   21   23   24   25   26   35   46   47   48   49   50   51   52   53
ring nodes :
    1   2   3   4   5   6   7   8   9   10   11   12   13   15   16   17   18   19   20   22   27
    28   29   30   31   32   33   34   36   37   38   39   40   41   42   43   44   45
chain bonds :
    1-21   9-14   19-22   19-23   23-24   23-25   25-26   34-35   37-46   38-48   39-49
    39-50   46-47   50-51   50-52   52-53
ring bonds :
    1-2   1-5   1-20   2-6   3-4   3-6   4-5   6-7   7-8   8-12   9-10   9-13   10-11
    10-18   11-12   11-15   12-13   13-19   15-16   16-17   17-18   19-20   22-27
    22-31   27-28   28-29   28-32   29-30   29-34   30-31   32-33   32-36   32-40
    33-34   33-39   36-37   36-42   37-38   37-45   38-39   40-41   41-42   42-43
    43-44   44-45
exact/norm bonds :
    1-2   1-5   1-20   2-6   3-4   3-6   4-5   6-7   7-8   8-12   9-10   9-13   11-12
    12-13   13-19   19-20   23-24   23-25   28-32   29-34   32-33   32-36   32-40
    33-34   33-39   36-37   36-42   37-38   37-45   38-39   38-48   39-49   40-41
    41-42   42-43   43-44   44-45   50-51   50-52
exact bonds :
    1-21   9-14   19-22   19-23   25-26   34-35   37-46   39-50   46-47   52-53
normalized bonds :
    10-11   10-18   11-15   15-16   16-17   17-18   22-27   22-31   27-28   28-29
    29-30   30-31

Match level :
    1:Atom   2:Atom   3:Atom   4:Atom   5:Atom   6:Atom   7:Atom   8:Atom   9:Atom
    10:Atom

11:Atom 1●Atom 13:Atom 14:CLASS   Atom 16:Atom 17:Atom
18:Atom 19:Atom 20:Atom 21:CLASS 22:Atom 23:CLASS 24:CLASS
25:CLASS 26:CLASS 27:Atom 28:Atom 29:Atom 30:Atom 31:Atom 32:Atom
33:Atom 34:Atom 35:CLASS 36:Atom 37:Atom 38:Atom 39:Atom 40:Atom
41:Atom 42:Atom 43:Atom 44:Atom 45:Atom 46:CLASS 47:CLASS 48:CLASS
49:CLASS 50:CLASS 51:CLASS 52:CLASS 53:CLASS